1/1

FIG. 1

```
     1                                                              61
P1     IleArgLysArgXaaAlaArgCysMetGlnLysAspGlyXaaLysAlaAspGlyIleAsp
+      gatcmggaaacgyttsgctcggtgcatgcagaaggacgggwtgaaggcggacgggattgac
-      nctagkcctttgcraawcgagccacgtacgtcttcctgcccsacttccgcctgccctaactg
M1   SerXaaSerValXaaXaaGluThrCysAlaSerProProXaaSerProProArgSerGlnAr
M2     IleXaaPheArgLysAlaArgHisMetCysPheSerProXaaPheAlaSerProIleSerS

     62                                                             121
P1     AspAspAspAspIleAlaMetLysAspGlyThrAlaAspValLeuGlyGlyAlaGluArg
+      gacgacgacgacattgcgatgaaagatgggaccgcygacgtccttggcggggcggagcgc
-      ctgctgctgctgtaacgctactttctaccctggcgrctgcaggaaccgccccgcctcgcg
M1   gArgArgArgCysGlnSerSerHisLeuSerArgXaaArgGlyGlnArgProProAlaAr
M2     erSerSerSerMet

     122                                                            181
P1     GluAsnGlnAspAspGluAspGluAspValTyrAlaArgIleArgPheLeuProGluArg
+      gagaaccaagacgacgaggacgaggacgtctacgcgcgcatccgtttccttcctgagcgg
-      ctcttggttctgctgctcctgctcctgcagatgcgcgcgtaggcaaaggaaggactcgcc
M1   gSerGlyLeuArgArgProArgProArgArgArgAlaCysGlyAsnGlyGluGlnAlaPr

     182                                                            241
P1     ValPheAspThrSerAlaLeuLeuIleLeuLysPheSerLeuAlaAspAlaAspSerAla
+      gtatttgacacctccgcattgctgatcctgaagttctcgcttgcagacgctgattcagcg
-      cataaactgtggaggcgtaacgactaggacttcaagagcgaacgtctgcgactaagtcgc
M1   oIleGlnCysArgArgMetAlaSerGlySerThrArgAlaGlnLeuArgGlnAsnLeuAl

     242                                                            301
P1     ProLeuArgArgThrCysPheGlyArgCysLysProHisGlySerAspHisArgGlnPhe
+      ccgcttcgtcgcacctgctttggacgctgcaaaccgcacggctcggaccatcgtcagttt
-      ggcgaagcagcgtggacgaaacctgcgacgtttggcgtgccgagcctggtagcagtcaaa
M1   aAlaGluAspCysArgSerGlnValSerCysValAlaArgSerProGlyAspAspThrGl

     302                                                            361
P1     ProAlaSerGluValAsnPheArgProArgTrpThrLeuLeuSerLeuLeuSerLeuPro
+      cctgcttcagaggtgaatttccgaccccgttggactttgctctctcttctctctctaccc
-      ggacgaagtctccacttaaaggctggggcaacctgaaacgagagagaagagagagatggg
M1   uGlnLysLeuProSerAsnGlyValGlyAsnSerLysAlaArgGluGluArgGluValAr

     362      371
P1     AspAspAsp
+      gacgacgatc
-      ctgctgctag
M1   gArgArgAsp
```